



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/645,304
Source: PIPE
Date Processed by STIC: 8-28-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/efb/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

101645.304

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 X Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



OIPE

RAW SEQUENCE LISTING

DATE: 08/28/2003

PATENT APPLICATION: US/10/645,304

TIME: 09:03:34

Input Set : E:\126481_1001prj.ST25.txt

Output Set: N:\CRF4\08282003\J645304.raw

3 <110> APPLICANT: Samuel , Stupp I.
 5 <120> TITLE OF INVENTION: CHARGED PEPTIDE-AMPHIPHILE SOLUTIONS & SELF ASSEMBLED
 PEPTIDE
 6 NANOFIBER NETWORKS FORMED THEREBY
 8 <130> FILE REFERENCE: 126481.1001
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/645,304
 11 <141> CURRENT FILING DATE: 2003-08-21
 13 <150> PRIOR APPLICATION NUMBER: 60/406,016
 14 <151> PRIOR FILING DATE: 2002-08-21
 16 <160> NUMBER OF SEQ ID NOS: 22
 18 <170> SOFTWARE: PatentIn version 3.2
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 7
 22 <212> TYPE: PRT
 23 <213> ORGANISM: Cystine with a 16 carbon alkyl chain attached
 25 <400> SEQUENCE: 1
 27 Cys Cys Cys Cys Gly Gly Gly
 28 1 5
 31 <210> SEQ ID NO: 2
 32 <211> LENGTH: 7
 33 <212> TYPE: PRT
 34 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
 36 <400> SEQUENCE: 2
 38 Ala Ala Ala Ala Gly Gly Gly
 39 1 5
 42 <210> SEQ ID NO: 3
 43 <211> LENGTH: 7
 44 <212> TYPE: PRT
 45 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
 47 <400> SEQUENCE: 3
 49 Ser Leu Ser Leu Gly Gly Gly
 50 1 5
 53 <210> SEQ ID NO: 4
 54 <211> LENGTH: 7
 55 <212> TYPE: PRT
 56 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
 58 <400> SEQUENCE: 4
 60 Cys Cys Cys Cys Gly Gly Gly
 61 1 5
 64 <210> SEQ ID NO: 5
 65 <211> LENGTH: 7
 66 <212> TYPE: PRT
 67 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
 69 <400> SEQUENCE: 5

Not a valid
 response. See
 Item 10 on
 error summary
 sheet.

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Input Set : E:\126481_1001prj.ST25.txt

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71 Ala Ala Ala Ala Gly Gly Gly
72 1 5
75 <210> SEQ ID NO: 6
76 <211> LENGTH: 7
77 <212> TYPE: PRT
78 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
80 <400> SEQUENCE: 6
82 Ser Leu Ser Leu Gly Gly Gly
83 1 5
86 <210> SEQ ID NO: 7
87 <211> LENGTH: 7
88 <212> TYPE: PRT
89 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
91 <400> SEQUENCE: 7
93 Cys Cys Cys Cys Gly Gly Gly
94 1 5
97 <210> SEQ ID NO: 8
98 <211> LENGTH: 7
99 <212> TYPE: PRT
100 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
102 <400> SEQUENCE: 8
104 Ala Ala Ala Ala Gly Gly Gly
105 1 5
108 <210> SEQ ID NO: 9
109 <211> LENGTH: 7
110 <212> TYPE: PRT
111 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
113 <400> SEQUENCE: 9
115 Ser Leu Ser Leu Gly Gly Gly
116 1 5
119 <210> SEQ ID NO: 10
120 <211> LENGTH: 7
121 <212> TYPE: PRT
122 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
124 <400> SEQUENCE: 10
126 Cys Cys Cys Cys Gly Gly Gly
127 1 5
130 <210> SEQ ID NO: 11
131 <211> LENGTH: 7
132 <212> TYPE: PRT
133 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
135 <400> SEQUENCE: 11
137 Ala Ala Ala Ala Gly Gly Gly
138 1 5
141 <210> SEQ ID NO: 12
142 <211> LENGTH: 7
143 <212> TYPE: PRT
144 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
146 <400> SEQUENCE: 12

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Input Set : E:\126481_1001prj.ST25.txt

Output Set: N:\CRF4\08282003\J645304.raw

148 Ser Leu Ser Leu Gly Gly Gly
149 1 5
152 <210> SEQ ID NO: 13
153 <211> LENGTH: 7
154 <212> TYPE: PRT
155 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
157 <400> SEQUENCE: 13
159 Cys Cys Cys Cys Gly Gly Gly
160 1 5
163 <210> SEQ ID NO: 14
164 <211> LENGTH: 7
165 <212> TYPE: PRT
166 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
168 <400> SEQUENCE: 14
170 Ala Ala Ala Ala Gly Gly Gly
171 1 5
174 <210> SEQ ID NO: 15
175 <211> LENGTH: 7
176 <212> TYPE: PRT
177 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
179 <400> SEQUENCE: 15
181 Ser Leu Ser Leu Gly Gly Gly
182 1 5
185 <210> SEQ ID NO: 16
186 <211> LENGTH: 7
187 <212> TYPE: PRT
188 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
190 <400> SEQUENCE: 16
192 Cys Cys Cys Cys Gly Gly Gly
193 1 5
196 <210> SEQ ID NO: 17
197 <211> LENGTH: 7
198 <212> TYPE: PRT
199 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
201 <400> SEQUENCE: 17
203 Ala Ala Ala Ala Gly Gly Gly
204 1 5
207 <210> SEQ ID NO: 18
208 <211> LENGTH: 7
209 <212> TYPE: PRT
210 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
212 <400> SEQUENCE: 18
214 Ser Leu Ser Leu Gly Gly Gly
215 1 5
218 <210> SEQ ID NO: 19
219 <211> LENGTH: 7
220 <212> TYPE: PRT
221 <213> ORGANISM: Cystein with a 16 carbon alkyl chain attached
223 <400> SEQUENCE: 19

RAW SEQUENCE LISTING

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Input Set : E:\126481_1001prj.ST25.txt

Output Set: N:\CRF4\08282003\J645304.raw

225 Cys Cys Cys Cys Gly Gly Gly
226 1 5
229 <210> SEQ ID NO: 20
230 <211> LENGTH: 7
231 <212> TYPE: PRT
232 <213> ORGANISM: Alanine with a 16 carbon alkyl chain attached
234 <400> SEQUENCE: 20
236 Ala Ala Ala Ala Gly Gly Gly
237 1 5
240 <210> SEQ ID NO: 21
241 <211> LENGTH: 7
242 <212> TYPE: PRT
243 <213> ORGANISM: Serine with a 16 carbon alkyl chain attached
245 <400> SEQUENCE: 21
247 Ser Leu Ser Leu Gly Gly Gly
248 1 5
251 <210> SEQ ID NO: 22
252 <211> LENGTH: 7
253 <212> TYPE: PRT
254 <213> ORGANISM: x is 2,3-diaminopropionic acid
257 <220> FEATURE:
258 <221> NAME/KEY: misc_feature
259 <222> LOCATION: (5)..(7)
260 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
262 <400> SEQUENCE: 22
W--> 264 Ser Leu Ser Leu Xaa Xaa Xaa
265 1 5

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Input Set : E:\126481_1001prj.ST25.txt
Output Set: N:\CRF4\08282003\J645304.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:22; Xaa Pos. 5,6,7

VERIFICATION SUMMARY

DATE: 08/28/2003

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TIME: 09:03:35

Input Set : E:\126481_1001prj.ST25.txt

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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:264 M:341 W: (46) "n" or "Xaa" used, for .SEQ ID#:22 after pos.:0